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General Information

XP-002386149

Filliary Access

Primary Accession # BQ055725

Accession #

BQ055725

Entry Name

EMBL:BQ055725

Molecule Type

linear mRNA

Sequence Length

1060

Entry Division

HUM

Entry Data Class

EST

Sequence Version

BQ055725.1

Creation Date

01-APR-2002

Modification Date

01-APR-2002

Description

Description

AGENCOURT_6796360 NIH_MGC_99 Homo sapiens cDNA clone IMAGE:5807946

5', mRNA sequence.

Keywords

EST .:

Organism

Homo sapiens (human)

Organism Classification Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Euarchontoglires, Primates, Haplorrhini, Catarrhini, Hominidae, Homo.

References

1. NCI-MGC;

National Institutes of Health, Mammalian Gene Collection (MGC)

http://mgc.nci.nih.gov/

Unpublished.

Position 1-1060

Database Cross-references

RZPD IMAGp958C192051.

IMAGp958C192051Q.

UNILIB 23347; 9630.

Features

Key Location Qualifier Value

source 1..1060

organism Homo sapiens

lab_host DH10B (phage-resistant)

moi_type mRNA

clone_lib NIH_MGC_99

clone IMAGE:5807946

tissue_type lymphoma, cell line

note Organ: lymph; Vector: pOTB7; Site_1: Xhol; Site_2:

EcoRt; cDNA made by oligo-dT priming. Directionally cloned into EcoRt/Xhol sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for

average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library.

db_xref taxon:9606

db_xref RZPD:IMAGp958C192051

db_xref RZPD:IMAGp958C192051Q

db_xref UNILIB:23347

Sequence

Characteristics

Length: 1060 BP, A Count:264, C Count:348, G Count:223, T Count:224, Others Count:1

Sequence

>embl BQ055725 BQ055725 AGENCOURT 6796360 NIH MGC 99 Homo sapiens cDNA cctgttggtctctacccacaggggatcccctggctgctccaccatcagatttgggacacc acccccccggccccaccagagggcatcagctatgcctagaaggggaccacaacagactc gacaggatecaceggttgggcccaaggcaggaggaagggcgcccccaaactcccagg acgcctgcagcacccccacgcgccgctctccgcctctggggagcatcctgccaccccc gacacacacaccccggctacatcccgccttctcacgcttggtcaggcgctctggagatgt cggagatccaggcttttcctaaagagtcaggattggaaggcggactcccaccgtttgctg agetecacatgacaacagcagacgacaggeegcactgateccacetggettacaggtget gtcacacaggtacgccttcttgtgaatcatcagtcaccatttgcccctcgcagcaaccct gtettggaggaaggteettgeegaeaeeegeeteteeeeaeeettaggeteeettgetee tteatteaaceattgtttacatgtegtteaggaetgetgaetaaacaettteaageacag tggaccettgaacaacgcaggttcaaactgtgcaggtccaccccacagggatgtttttcaa tragagtratarragtgreertgretertretgretreertrecartrecetratgtett cotototgccaccotgaggcaacaagaaccagccotcotcotcagccactcaacatgaa gaacaccaggatggagacttctattgataatccacttcccctttagnaaaagctttccct taaggaaatteettaatggacattttaattteeceaagettagtttaaatggtaaaaata ccggaataaataacccttccaaaaaaaaaaaaaaaaactgaaggcatttaattctcggg ggccgaaaaaagaaggtaatggaaagggcacctggggccctaactggtttccctggaccc aactttttcctggtaacaaaaagggagagaatcttcccgg

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